OIPE O2 co

Serial Number: _09/369,236_

CRF Processing Dat : _8/24/99_
Edited by: _____
V rified by: ___/C___ (STIC staff)

☐ Changed a file from non-ASCII to ASCII

☐ Changed the margins in cases where the sequence text was "wrapped" down to the next line.

☐ Edited a format error in the Current Application Data section, specifically:
_____

☐ Edited the Current Application Data section with the actual current number. The number inputted by the applicant was ☐ the prior application data; or ☐ other _____.

ENTERED

☐ Added the mandatory heading and subheadings for "Current Application Data".

☐ Edited the "Number of Sequences" field. The applicant spelled out a number instead of using an integer.

☐ Changed the spelling of a mandatory field (the headings or subheadings), specifically:
_____

☐ Corrected the SEQ ID NO when obviously incorrect. The sequence numbers that were edited were:
_____

☐ Inserted or corrected a nucleic number at the end of a nucleic line. SEQ ID NO's edited:
_____

☑ Corrected subheading placement. All responses must be on the same line as each subheading. If the applicant placed a response below the subheading, this was moved to its appropriate place.

☐ Inserted colons after headings/subheadings. Headings edited included:
_____

☑ Deleted extra, invalid, headings used by an applicant, specifically:
_(A) NAmE:_____

☐ Deleted: ☐ non-ASCII "garbage" at the beginning/end of files; ☐ secretary initials/filename at end of file; ☐ page numbers throughout text; ☐ other invalid text, such as_____ .

☐ Inserted mandatory headings, specifically: _____

☐ Corrected an obvious error in the response, specifically:
_____

☐ Edited identifiers where upper case is used but lower case is required, or vice versa.

☐ Corrected an error in the Number of Sequences field, specifically:
_____

☐ A "Hard Page Break" code was inserted by the applicant. All occurrences had to be deleted.

☐ Deleted *ending* stop codon in amino acid sequences and adjusted the "(A)Length:" field accordingly (error due to a PatentIn bug). Sequences corrected: _____

☐ Other:
_____
_____
_____

*Examiner: Th above corrections must be communicated to the applicant in th first Office Action. DO NOT send a copy of this form.

3/1/95

OIPE

PAGE: 1

RAW SEQUENCE LISTING
PATENT APPLICATION *US/09/369,236*

DATE: 08/24/1999
TIME: 14:38:53

INPUT SET: S33041.raw

This Raw Listing contains the General
Information Section and up to the first 5 pages.

```
 1                              SEQUENCE LISTING
 2                                                        Does Not Comply
 3      (1)     General Information:                  Corrected Diskette Needed
 4
 5          (i) APPLICANT: T
 6              (A) NAME: Grant A. Krafft, William L. Klein, Brett A. Chromy,
 7      Mary P. Lambert, Caleb E. Finch, Todd Morgan, Pat Wals,
 8      Irina Rozovsky, Ann Barlow
 9
10          (ii) TITLE OF INVENTION: Amyloid ( Protein (Globular Assembly
11      and Uses Thereof)
12
13          (iii) NUMBER OF SEQUENCES: 4
14
15          (iv) CORRESPONDENCE ADDRESS:
16          (A) ADDRESSEE:  McDonnell Boehnen Hulbert & Berghoff
17          (B) STREET:   300 South Wacker Drive
18          (C) CITY:   Chicago
19              (D) STATE:   IL
20          (E) COUNTRY:  USA
21          (F) ZIP:  60606
22
23          (v) COMPUTER READABLE FORM:
24              (A) MEDIUM TYPE: Floppy disk
25              (B) COMPUTER: IBM PC compatible
26              (C) OPERATING SYSTEM: PC-DOS/MS-DOS
27              (D) SOFTWARE: PatentIn Release #1.0, Version #1.30 (US)
28
29          (vi) CURRENT APPLICATION DATA:
30      (A) APPLICATION NUMBER: unassigned
31      (B) FILING DATE: 04-AUG-1999
32      (C) CLASSIFICATION DATA:
33
34          (vii) PRIOR APPLICATION DATA:
35              (A) APPLICATION NUMBER: US 08/796,089
36              (B) FILING DATE: 05-FEB-1997
37
38
39      (2) INFORMATION FOR SEQ ID NO: 1:
40
41          (i) SEQUENCE CHARACTERISTICS:
42              (A) LENGTH: 20 base pairs
43              (B) TYPE: nucleic acid
44              (C) STRANDEDNESS: single
45              (D) TOPOLOGY: linear
46
```

INPUT SET: S33041.raw

```
47        (ii) MOLECULE TYPE: other nucleic acid
48
49
50        (xi) SEQUENCE DESCRIPTION: SEQ ID NO: 1:
51
52        GCACCTTCTT TCCCTTCATC 20
53
54   (2) INFORMATION FOR SEQ ID NO: 2:
55
56        (i) SEQUENCE CHARACTERISTICS:
57             (A) LENGTH: 20 base pairs
58             (B) TYPE: nucleic acid
59             (C) STRANDEDNESS: single
60             (D) TOPOLOGY: linear
61
62        (ii) MOLECULE TYPE: other nucleic acid
63
64
65        (xi) SEQUENCE DESCRIPTION: SEQ ID NO: 2:
66
67        TGCTGATGTA CCAGTTGGGG 20
68
69
70   (2) INFORMATION FOR SEQ ID NO: 3:
71
72        (i) SEQUENCE CHARACTERISTICS:
73             (A) LENGTH: 19 base pairs
74             (B) TYPE: nucleic acid
75             (C) STRANDEDNESS: single
76             (D) TOPOLOGY: linear
77
78        (ii) MOLECULE TYPE: other nucleic acid
79
80
81        (xi) SEQUENCE DESCRIPTION: SEQ ID NO: 3:
82
83        CAGTCCTTGA CCTGCGACC 19
84
85
86   (2) INFORMATION FOR SEQ ID NO: 4:
87
88        (i) SEQUENCE CHARACTERISTICS:
89             (A) LENGTH: 19 base pairs
90       (B) TYPE: nucleic acid
91             (C) STRANDEDNESS: single
92             (D) TOPOLOGY: linear
93
94        (ii) MOLECULE TYPE: other nucleic acid
95
96        (xi) SEQUENCE DESCRIPTION: SEQ ID NO: 4:
97
98        GCCTCACATC ACATCCTTG 19
99
```

SEQUENCE VERIFICATION REPORT
PATENT APPLICATION   *US/09/369,236*

INPUT SET: S33041.raw

Line	Error	Original Text
5	Mandatory Value Not Present	(i) APPLICANT:
6	Unknown or Misplaced Identifier	(A) NAME: Grant A. Krafft, William L. Klein, Brett A. Ch
30	Wrong application Serial Number	(A) APPLICATION NUMBER: unassigned